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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/468,647

DATE: 07/24/2000  
TIME: 13:42:13

Input Set : A:\pto.txt  
Output Set: N:\CRF3\07242000\I468647.raw

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3 <110> APPLICANT: Gordon, Robert D.  
4 Sprengel, Jorg J.  
5 Yon, Jeffrey R.  
6 Dijkmans, Josiena J.H.  
7 Gosiewska, Anna  
8 Dhanaraj, Sridevi N.  
9 Xu, Jean  
11 <120> TITLE OF INVENTION: Vascular Endothelial Growth Factor-X  
13 <130> FILE REFERENCE: B0192/7011  
15 <140> CURRENT APPLICATION NUMBER: US 09/468,647  
16 <141> CURRENT FILING DATE: 1999-12-21  
18 <150> PRIOR APPLICATION NUMBER: GB 9828377.3  
19 <151> PRIOR FILING DATE: 1998-12-22  
21 <150> PRIOR APPLICATION NUMBER: US 60/124,967  
22 <151> PRIOR FILING DATE: 1999-03-18  
24 <150> PRIOR APPLICATION NUMBER: US 60/164,131  
25 <151> PRIOR FILING DATE: 1999-11-08  
27 <160> NUMBER OF SEQ ID NOS: 29  
29 <170> SOFTWARE: PatentIn Ver. 2.0  
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32 <211> LENGTH: 323  
33 <212> TYPE: PRT  
34 <213> ORGANISM: Homo sapiens  
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38 1 5 10 15  
40 Tyr Gly Val Gln Asp Pro Gln His Glu Arg Ile Ile Thr Val Ser Thr  
41 20 25 30  
43 Asn Gly Ser Ile His Ser Pro Arg Phe Pro His Thr Tyr Pro Arg Asn  
44 35 40 45  
46 Thr Val Leu Val Trp Arg Leu Val Ala Val Glu Glu Asn Val Trp Ile  
47 50 55 60  
49 Gln Leu Thr Phe Asp Glu Arg Phe Gly Leu Glu Asp Pro Glu Asp Asp  
50 65 70 75 80  
52 Ile Cys Lys Tyr Asp Phe Val Glu Val Glu Glu Pro Ser Asp Gly Thr  
53 85 90 95  
55 Ile Leu Gly Arg Trp Cys Gly Ser Gly Thr Val Pro Gly Lys Gln Ile  
56 100 105 110  
58 Ser Lys Gly Asn Gln Ile Arg Ile Arg Phe Val Ser Asp Glu Tyr Phe  
59 115 120 125  
61 Pro Ser Glu Pro Gly Phe Cys Ile His Tyr Asn Ile Val Met Pro Gln  
62 130 135 140  
64 Phe Thr Glu Ala Val Ser Pro Ser Val Leu Pro Pro Ser Ala Leu Pro  
65 145 150 155 160  
67 Leu Asp Leu Leu Asn Ala Ile Thr Ala Phe Ser Thr Leu Glu Asp  
68 165 170 175  
70 Leu Ile Arg Tyr Leu Glu Pro Glu Arg Trp Gln Leu Asp Leu Glu Asp

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73 Leu Tyr Arg Pro Thr Trp Gln Leu Leu Gly Lys Ala Phe Val Phe Gly
74          195          200          205
76 Arg Lys Ser Arg Val Val Asp Leu Asn Leu Leu Thr Glu Glu Val Arg
77          210          215          220
79 Leu Tyr Ser Cys Thr Pro Arg Asn Phe Ser Val Ser Ile Arg Glu Glu
80 225          230          235          240
82 Leu Lys Arg Thr Asp Thr Ile Phe Trp Pro Gly Cys Leu Leu Val Lys
83          245          250          255
85 Arg Cys Gly Gly Asn Cys Ala Cys Cys Leu His Asn Cys Asn Glu Cys
86          260          265          270
88 Gln Cys Val Pro Ser Lys Val Thr Lys Lys Tyr His Glu Val Leu Gln
89          275          280          285
91 Leu Arg Pro Lys Thr Gly Val Arg Gly Leu His Lys Ser Leu Thr Asp
92          290          295          300
94 Val Ala Leu Glu His His Glu Glu Cys Asp Cys Val Cys Arg Gly Ser
95 305          310          315          320
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111          20          25          30
113 Ser Ser Asn Lys Glu Gln Tyr Gly Val Gln Asp Pro Gln His Glu Arg
114          35          40          45
116 Ile Ile Thr Val Ser Thr Asn Gly Ser Ile His Ser Pro Arg Phe Pro
117          50          55          60
119 His Thr Tyr Pro Arg Asn Thr Val Leu Val Trp Arg Leu Val Ala Val
120 65          70          75          80
122 Glu Glu Asn Val Trp Ile Gln Leu Thr Phe Asp Glu Arg Phe Gly Leu
123          85          90          95
125 Glu Asp Pro Glu Asp Asp Ile Cys Lys Tyr Asp Phe Val Glu Val Glu
126          100          105          110
128 Glu Pro Ser Asp Gly Thr Ile Leu Gly Arg Trp Cys Gly Ser Gly Thr
129          115          120          125
131 Val Pro Gly Lys Gln Ile Ser Lys Gly Asn Gln Ile Arg Ile Arg Phe
132          130          135          140
134 Val Ser Asp Glu Tyr Phe Pro Ser Glu Pro Gly Phe Cys Ile His Tyr
135 145          150          155          160
137 Asn Ile Val Met Pro Gln Phe Thr Glu Ala Val Ser Pro Ser Val Leu
138          165          170          175
140 Pro Pro Ser Ala Leu Pro Leu Asp Leu Leu Asn Asn Ala Ile Thr Ala
141          180          185          190
143 Phe Ser Thr Leu Glu Asp Leu Ile Arg Tyr Leu Glu Pro Glu Arg Trp
144          195          200          205
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146 Gln Leu Asp Leu Glu Asp Leu Tyr Arg Pro Thr Trp Gln Leu Leu Gly  
147 210 215 220  
149 Lys Ala Phe Val Phe Gly Arg Lys Ser Arg Val Val Asp Leu Asn Leu  
150 225 230 235 240  
152 Leu Thr Glu Glu Val Arg Leu Tyr Ser Cys Thr Pro Arg Asn Phe Ser  
153 245 250 255  
155 Val Ser Ile Arg Glu Glu Leu Lys Arg Thr Asp Thr Ile Phe Trp Pro  
156 260 265 270  
158 Gly Cys Leu Leu Val Lys Arg Cys Gly Gly Asn Cys Ala Cys Cys Leu  
159 275 280 285  
161 His Asn Cys Asn Glu Cys Gln Cys Val Pro Ser Lys Val Thr Lys Lys  
162 290 295 300  
164 Tyr His Glu Val Leu Gln Leu Arg Pro Lys Thr Gly Val Arg Gly Leu  
165 305 310 315 320  
167 His Lys Ser Leu Thr Asp Val Ala Leu Glu His His Glu Glu Cys Asp  
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182 gtacaagatc ctcagcatga gagaattatt actgtgtcta ctaatggaag tattcacagc 180  
183 ccaagggttc ctcatactta tccaagaaat acggctcttg tatggagatt agtagcagta 240  
184 gaggaaaaatg tatggataca acttacgttt gatgaaagat ttgggcttga agaccagaa 300  
185 gatgacatat gcaagtatga tttttagtaa gttgaggaac ccagtgtatg aactatatta 360  
186 gggcgctggt gtggttcttg tactgtacca ggaacacaga tttctaaagg aaatcaaatt 420  
187 aggataagat ttgtatctga tgaatatttt ccttctgaac cagggttctg catccactac 480  
188 aacattgtca tgcacaatt cacagaagct gtgagtcctt cagtgtctacc cccttcagct 540  
189 ttgccactgg acctgcttaa taatgctata actgccttta gtaccttggg agaccttatt 600  
190 cgatatcttg aaccagagag atggcagttg gacttagaag atctatatag gccaaacttg 660  
191 caacttcttg gcaaggcttt tgtttttgga agaaaatcca gagtgggtgga tctgaacctt 720  
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193 gaagaaactaa agagaaccga taccattttc tggccagggt gtctcctggt taaacgctgt 840  
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195 gttactaaaa aataccacga ggtccttcag ttgagaccaa agaccgggtg caggggattg 960  
196 cacaatcac tcaccgacgt ggccttgagg caccatgagg agtgtgactg tgtgtgcaga 1020  
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202 <213> ORGANISM: Artificial Sequence  
204 <220> FEATURE:  
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208 aaaatgtatg gatacaactt ac

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223 <212> TYPE: DNA  
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230 ttctctaaagg aaatcaaatt ag  
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241 gataagattt gtatctgatg  
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244 <211> LENGTH: 17  
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256 <212> TYPE: DNA  
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284 <400> SEQUENCE: 11 20  
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VERIFICATION SUMMARY  
PATENT APPLICATION: US/09/468,647

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